HYS-38CIP sequence listing
SEQUENCE LISTING

<110> Dedera, Douglas
Yamazaki, Victoria
Asundi, Vinod
Liu, Chenghua
Tang, Y. Tom
Drmanac, Radoje T.

<120> Methods of Therapy and Diagnosis Using Insulin-like Growth Factor Binding Protein-like Polypeptides and Polynucleotides

<130> HYS-38CIP

<140> Not Yet Assigned
<141> 2002-02-27

<150> 09/784,748
<151> 2001-02-14

<150> 09/649,167
<151> 2000-08-23

<150> 09/540,217
<151> 2000-03-31

<160> 14

<170> PatentIn version 3.1

<210> 1
<211> 375

<212> DNA

<213> Homo sapiens

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tcaccggggc gcaggtgggc ctgtcctgtg aagtgagggc tgtgcctacc ccagtcatca 180
cgtggagaaa ggtcacgaag tcccctgagg gcacccaagc actggaggag ctgcctgggg 240
accatgtcaa tatagctgtc caagtgcgag ggggcccttc tgaccatgag gccacggcct 300
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gaagagtagg gacatactta gatgagcagt agaatccctg gtatattctg cacacatctc 300
cctctgagct tcttagcatg caaagacaag agctgtgaac atgaaggtgt gtccatgaga 360
tgaaaagacc agttgtgttt tggggctgga gggaatattt cctctgtatt cttttagaaa 420
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<212> DNA

<213> Homo sapiens

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tcaccggggc gcaggtgggc ctgtcctgtg aagtgagggc tgtgcctacc ccagtcatca 180

cgtggagaaa ggtcacgaag tcccctgagg gcacccaagc actggaggag ctgcctgggg 240

accatgtcaa tatagctgtc caagtgcgag ggggcccttc tgaccatgag gccacggcct 300

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<211> 1250

<212> DNA

<213> Homo sapiens

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Met Pro Arg Leu Ser Leu Leu Leu Pro Leu Leu
1 5 10

ctt ctg ctg ctg ctg ccg ctg ctg ccg ccg ctg tcc ccg agc ctc ggg 159
Leu Leu Leu Leu Leu Pro Leu Leu Pro Pro Leu Ser Pro Ser Leu Gly
15 20 25

atc cgc gac gtg ggc ggc cgg cgc ccc aag tgt ggt ccg tgc cgg cca 207
Ile Arg Asp Val Gly Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro
30 35 40

gag ggc tgc ccg gcg cct gcg ccc tgc ccg gcg ccc ggg atc tcg gcg 255
Glu Gly Cys Pro Ala Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala
45 50 55

ctc gac gag tgc ggc tgc tgc gcc cgc tgc ctg gga gcc gag ggc gcg 303
Leu Asp Glu Cys Gly Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala
60 65 70 75

agc tgc ggg ggc cgc gcc ggc ggg cgc tgt ggc ccc ggc ctg gta tgc 351
Ser Cys Gly Gly Arg Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys
80 85 90

gcg agc cag gcc gct ggg gca gcg ccc gag ggc acc ggg ctc tgc gtg 399
Ala Ser Gln Ala Ala Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val
95 100 105

tgc gcg cag cgc ggc acc gtc tgc ggc tcc gac ggt cgc tcg tac ccc 447
Cys Ala Gln Arg Gly Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro
110 115 120

agc gtc tgc gcg ctg cgc ctg cgc gct cgg cac acg ccc cgc gcg cac 495
Ser Val Cys Ala Leu Arg Leu Arg Ala Arg His Thr Pro Arg Ala His
125 130 135

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ccc ggt cac ctg cac aag gcg cgc gac ggc cct tgc gag ttc gct cct      543
Pro Gly His Leu His Lys Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro
140                 145                 150                 155

gtg gtc gtc gtt cct ccc cga agt gtt cac aac gtc acc ggg gcg cag      591
Val Val Val Val Pro Pro Arg Ser Val His Asn Val Thr Gly Ala Gln
                160                 165                 170

gtg ggc ctg tcc tgt gaa gtg agg gct gtg cct acc cca gtc atc acg      639
Val Gly Leu Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr
            175                 180                 185

tgg aga aag gtc acg aag tcc cct gag ggc acc caa gca ctg gag gag      687
Trp Arg Lys Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu
        190                 195                 200

ctg cct ggg gac cat gtc aat ata gct gtc caa gtg cga ggg ggc cct      735
Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro
    205                 210                 215

tct gac cat gag gcc acg gcc tgg att ttg atc aac ccc ctg cga aag      783
Ser Asp His Glu Ala Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys
220                 225                 230                 235

gag gat gag ggt gtg tac cag tgc cat gca gcc aac atg gtg gga gag      831
Glu Asp Glu Gly Val Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu
                240                 245                 250

gct gag tcc cac agc aca gtg acg gtt cta gat ctg agt aaa tac agg      879
Ala Glu Ser His Ser Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg
            255                 260                 265

agc ttc cac ttc cca gct ccc gat gac cgc atg tga tggagaaatg           925
Ser Phe His Phe Pro Ala Pro Asp Asp Arg Met
        270                 275

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<210> 6

<211> 278

<212> PRT

<213> Homo sapiens

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            20                  25                  30

Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro Glu Gly Cys Pro Ala
        35                  40                  45
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Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala Leu Asp Glu Cys Gly
    50                  55                  60

Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala Ser Cys Gly Gly Arg
65                  70                  75                  80

Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys Ala Ser Gln Ala Ala
                85                  90                  95

Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val Cys Ala Gln Arg Gly
            100                 105                 110

Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro Ser Val Cys Ala Leu
        115                 120                 125

Arg Leu Arg Ala Arg His Thr Pro Arg Ala His Pro Gly His Leu His
    130                 135                 140

Lys Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro Val Val Val Val Pro
145                 150                 155                 160

Pro Arg Ser Val His Asn Val Thr Gly Ala Gln Val Gly Leu Ser Cys
                165                 170                 175

Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys Val Thr
            180                 185                 190

Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly Asp His
        195                 200                 205

Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro Ser Asp His Glu Ala
    210                 215                 220

Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys Glu Asp Glu Gly Val
225                 230                 235                 240

Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu Ala Glu Ser His Ser
                245                 250                 255

Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg Ser Phe His Phe Pro
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Ala Pro Asp Asp Arg Met
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<210> 7

<211> 837

<212> DNA

<213> Homo sapiens

<400> 7

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tgccggccag agggctgccc ggcgcctgcg ccctgcccgg cgcccgggat ctcggcgctc 180

gacgagtgcg gctgctgcgc ccgctgcctg ggagccgagg gcgcgagctg cgggggccgc 240

gccggcgggc gctgtggccc cggcctggta tgcgcgagcc aggccgctgg ggcagcgccc 300

gagggcaccg ggctctgcgt gtgcgcgcag cgcggcaccg tctgcggctc cgacggtcgc 360

tcgtacccca gcgtctgcgc gctgcgcctg cgcgctcggc acacgccccg cgcgcacccc 420

ggtcacctgc acaaggcgcg cgacggccct tgcgagttcg ctcctgtggt cgtcgttcct 480

ccccgaagtg ttcacaacgt caccggggcg caggtgggcc tgtcctgtga agtgagggct 540

gtgcctaccc cagtcatcac gtggagaaag gtcacgaagt cccctgaggg cacccaagca 600

ctggaggagc tgcctgggga ccatgtcaat atagctgtcc aagtgcgagg gggcccttct 660

gaccatgagg ccacggcctg gattttgatc aacccccctgc gaaaggagga tgagggtgtg 720

taccagtgcc atgcagccaa catggtggga gaggctgagt cccacagcac agtgacggtt 780

ctagatctga gtaaatacag gagcttccac ttcccagctc ccgatgaccg catgtga 837

<210> 8

<211> 16

<212> PRT

<213> Homo sapiens

<400> 8

Asp Glu Cys Gly Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala Ser
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<210> 9

<211> 27

<212> PRT

<213> Homo sapiens

HYS-38CIP sequence listing

<400> 9

Met Pro Arg Leu Ser Leu Leu Leu Pro Leu Leu Leu Leu Leu Leu Leu
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Pro Leu Leu Pro Pro Leu Ser Pro Ser Leu Gly
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<210> 10

<211> 251

<212> PRT

<213> Homo sapiens

<400> 10

Ile Arg Asp Val Gly Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro
1 5 10 15

Glu Gly Cys Pro Ala Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala
20 25 30

Leu Asp Glu Cys Gly Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala
35 40 45

Ser Cys Gly Gly Arg Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys
50 55 60

Ala Ser Gln Ala Ala Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val
65 70 75 80

Cys Ala Gln Arg Gly Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro
85 90 95

Ser Val Cys Ala Leu Arg Leu Arg Ala Arg His Thr Pro Arg Ala His
100 105 110

Pro Gly His Leu His Lys Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro
115 120 125

Val Val Val Val Pro Pro Arg Ser Val His Asn Val Thr Gly Ala Gln
130 135 140

Val Gly Leu Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr
145 150 155 160

HYS-38CIP sequence listing

Trp Arg Lys Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu
165 170 175

Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro
180 185 190

Ser Asp His Glu Ala Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys
195 200 205

Glu Asp Glu Gly Val Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu
210 215 220

Ala Glu Ser His Ser Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg
225 230 235 240

Ser Phe His Phe Pro Ala Pro Asp Asp Arg Met
245 250

<210> 11

<211> 103

<212> PRT

<213> Homo sapiens

<400> 11

Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro Val Val Val Val Pro Pro
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Arg Ser Val His Asn Val Thr Gly Ala Gln Val Gly Leu Ser Cys Glu
20 25 30

Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys Val Thr Lys
35 40 45

Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly Asp His Val
50 55 60

Asn Ile Ala Val Gln Val Arg Gly Gly Pro Ser Asp His Glu Ala Thr
65 70 75 80

Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys Glu Asp Glu Gly Val Tyr
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Gln Cys His Ala Ala Asn Met
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HYS-38CIP sequence listing

<210> 12

<211> 390

<212> PRT

<213> Homo sapiens

<400> 12

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            20                  25                  30

Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro Glu Gly Cys Pro Ala
        35                  40                  45

Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala Leu Asp Glu Cys Gly
    50                  55                  60

Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala Ser Cys Gly Gly Arg
65                  70                  75                  80

Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys Ala Ser Gln Ala Ala
                85                  90                  95

Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val Cys Ala Gln Arg Gly
            100                 105                 110

Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro Ser Val Cys Ala Leu
        115                 120                 125

Arg Leu Arg Ala Arg His Thr Pro Arg Ala His Pro Gly His Leu His
    130                 135                 140

Lys Ala Arg Asp Gly Pro Cys Glu Phe Val Pro Ile Thr Arg Phe Tyr
145                 150                 155                 160

Asn Cys Phe Pro Gln Pro Leu Ile His Arg Gln Phe Ser Leu Ser Pro
                165                 170                 175

Asp Arg Arg Gln Ser Glu Thr Leu Ser Lys Lys Lys Lys Lys Lys Glu
            180                 185                 190

Glu Glu Glu Glu Glu Glu Glu Glu Gly Glu Glu Glu Lys Glu Glu Glu
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195 200 205

Gly Cys Lys Ser Asn Phe Gln His Thr Ile Asn Phe Lys Glu Ile Ser
210 215 220

Glu Gly Phe Gly Lys Ile Phe Ser Phe Gln Pro Ser Met Ile Asp Ile
225 230 235 240

Ile Asp Glu Ala Ser Thr Leu His Val Ala Gln His Ala Val Val Leu
245 250 255

Asp Ala Arg Val Ala Glu Leu Leu Ser Asn Ala Ala Pro Val Val Val
260 265 270

Val Pro Pro Arg Ser Val His Asn Val Thr Gly Ala Gln Val Gly Leu
275 280 285

Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys
290 295 300

Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly
305 310 315 320

Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro Ser Asp His
325 330 335

Glu Ala Thr Ala Trp Ile Leu Val Ser Asp Leu His His Cys Leu Lys
340 345 350

Ala Leu Pro Thr Tyr Ser Tyr Ser Ser Thr Leu Ser Pro Ser Gln Val
355 360 365

Phe Leu Leu Ile His Leu Leu His Ile Gly Pro Tyr Pro Gly Ala Cys
370 375 380

Ile Leu Glu Ala Pro Pro
385 390

<210> 13

<211> 268

<212> PRT

<213> Mus musculus

<400> 13

HYS-38CIP sequence listing

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Gly Leu Gly Leu Arg Asp Ala Gly Arg Arg His Pro Glu Cys Ser Pro
            20                  25                  30

Cys Gln Gln Asp Arg Cys Pro Ala Pro Ser Pro Cys Pro Ala Pro Trp
        35                  40                  45

Ile Ser Ala Arg Asp Glu Cys Gly Cys Cys Ala Arg Cys Leu Gly Ala
    50                  55                  60

Glu Gly Ala Ser Cys Gly Gly Pro Val Gly Ser Arg Cys Gly Pro Gly
65                  70                  75                  80

Leu Val Cys Ala Ser Arg Ala Ser Gly Thr Ala Pro Glu Gly Thr Gly
                85                  90                  95

Leu Cys Val Cys Ala Gln Arg Gly Ala Val Cys Gly Ser Asp Gly Arg
            100                 105                 110

Ser Tyr Ser Ser Ile Cys Ala Leu Arg Leu Arg Ala Arg His Ala Pro
        115                 120                 125

Arg Ala His His Gly His Leu His Lys Ala Arg Asp Gly Pro Cys Glu
    130                 135                 140

Phe Ala Pro Val Val Leu Met Pro Pro Arg Asp Ile His Asn Val Thr
145                 150                 155                 160

Gly Thr Gln Val Phe Leu Ser Cys Glu Val Lys Ala Val Pro Thr Pro
                165                 170                 175

Val Ile Thr Trp Lys Lys Val Lys His Ser Pro Glu Gly Thr Glu Gly
            180                 185                 190

Leu Glu Glu Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg
        195                 200                 205

Gly Gly Pro Ser Asp His Glu Thr Thr Ser Trp Ile Leu Ile Asn Pro
    210                 215                 220

Leu Arg Lys Glu Asp Glu Gly Val Tyr His Cys His Ala Ala Asn Ala
225                 230                 235                 240

Ile Gly Glu Ala Gln Ser His Gly Thr Val Thr Val Leu Asp Leu Asn
                245                 250                 255
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Arg Tyr Lys Ser Leu Tyr Ser Ser Val Pro Gly Asp
260 265

<210> 14

<211> 264

<212> PRT

<213> Homo sapiens

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Pro Ser Leu Arg Ala Leu Leu Leu Gly Ala Ala Gly Leu Leu Leu Leu
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Leu Leu Pro Leu Ser Ser Ser Ser Ser Ser Asp Thr Cys Gly Pro Cys
20 25 30

Glu Pro Ala Ser Cys Pro Pro Leu Pro Pro Leu Gly Cys Leu Leu Gly
35 40 45

Glu Thr Arg Asp Ala Cys Gly Cys Cys Pro Met Cys Ala Arg Gly Glu
50 55 60

Gly Glu Pro Cys Gly Gly Gly Gly Ala Gly Arg Gly Tyr Cys Ala Pro
65 70 75 80

Gly Met Glu Cys Val Lys Ser Arg Lys Arg Arg Lys Gly Lys Ala Gly
85 90 95

Ala Ala Ala Gly Gly Pro Gly Val Ser Gly Val Cys Val Cys Lys Ser
100 105 110

Arg Tyr Pro Val Cys Gly Ser Asp Gly Thr Thr Tyr Pro Ser Gly Cys
115 120 125

Gln Leu Arg Ala Ala Ser Gln Arg Ala Glu Ser Arg Gly Glu Lys Ala
130 135 140

Ile Thr Gln Val Ser Lys Gly Thr Cys Glu Gln Gly Pro Ser Ile Val
145 150 155 160

Thr Pro Pro Lys Asp Ile Trp Asn Val Thr Gly Ala Gln Val Tyr Leu
165 170 175

Ser Cys Glu Val Ile Gly Ile Pro Thr Pro Val Leu Ile Trp Asn Lys
180 185 190

HYS-38CIP sequence listing

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Val Lys Arg Gly His Tyr Gly Val Gln Arg Thr Glu Leu Leu Pro Gly
        195                 200                 205

Asp Arg Asp Asn Leu Ala Ile Gln Thr Arg Gly Gly Pro Glu Lys His
    210                 215                 220

Glu Val Thr Gly Trp Val Leu Val Ser Pro Leu Ser Lys Glu Asp Ala
225                 230                 235                 240

Gly Glu Tyr Glu Cys His Ala Ser Asn Phe Gln Gly Gln Ala Ser Ala
                245                 250                 255

Ser Ala Lys Ile Thr Val Val Asp
            260
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